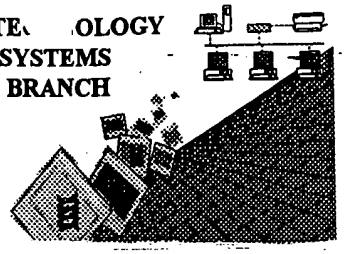
0590
0618

BIOTE OLOGY
SYSTEMS
BRANCH



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/602,362C
Source: OIPE
Date Processed by STIC: 6/17/02

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

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PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO **REDUCE** ERRORED SEQUENCE LISTINGS, **PLEASE** USE THE **CHECKER VERSION 3.1 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
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Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<**http://www.uspto.gov/ebc/efs/downloads/documents.htm**>, **EFS Submission User Manual** - ePAVE)
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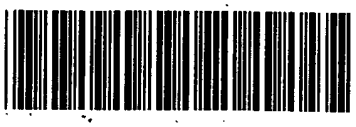
Revised 01/29/2002



OIPE

Does Not Comply
Corrected Diskette Needed

Errors on pp. 1 + 5

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/602,362C

DATE: 06/17/2002
TIME: 13:47:25

Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF3\06172002\I602362C.raw

1 <110> APPLICANT: Jager, Dirk
2 Scanlan, Matthew
3 Gure, Ali
4 Jager, Elke
5 Knuth, Alexander
6 Old, Lloyd
7 Chen, Yao-tseng
9 <120> TITLE OF INVENTION: Isolated Nucleic Acid Molecules Encoding Cancer Associated Antigens,
10 the Antigens per se, and Uses Thereof
12 <130> FILE REFERENCE: LUD 5615.1
C--> 14 **<140> CURRENT APPLICATION NUMBER: US/09/602,362C**
C--> 16 **<141> CURRENT FILING DATE: 2002-06-10**
18 <150> PRIOR APPLICATION NUMBER: 09/451,739
20 <151> PRIOR FILING DATE: 1999-11-30
22 <160> NUMBER OF SEQ ID NOS: 29
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 1533
27 <212> TYPE: DNA
28 <213> ORGANISM: Homo sapiens
W--> 29 **<220> FEATURE:**
30 <221> NAME/KEY: CDS
31 <222> LOCATION: 235
W--> 32 **<400> SEQUENCE: 1**

must explain what residue n represents, see p. 6

33 ggttttccac gttggacaag tgcggctcgg cggccagcgg agcgcgcccc ttcccgctgc 60
35 ccgctccgct cctctcttct acccagccca gtgggcgagt gggcagcggc ggccgcggcg 120
37 ctgggccctc tcccgccggt gtgtgcgcgc tcgtacgcgc ggcccccggc gccagccccg 180
W--> 39 **ccgcctgaga gggggcctgc gccgccggcc ggggcgtgcg cccgggagcc accgncaccg 240**
41 cggcccgcgc cctcaggcgc tggggtcccc gcggacccgg aggcggcgga cgggctcggc 300
43 agatgtagcc gccgggccga agcaggagcc ggcgggggcg cgccgggaga gcgagggctt 360
45 tgcattttgc agtgctattt tttgaggggg gcggagggtg gaggaagtcg gaaagccgcg 420
47 ccgagtcgcc ggggacctcc ggggtgaacc atgttgagtc ctgccaacgg ggagcagctc 480
49 cacctggtga actatgtgga ggactacctg gactccatcg agtccctgcc tttcgacttg 540
51 cagagaaatg tctcgctgat gcgggagatc gacgcgaaat accaagagat cctgaaggag 600
53 ctagacgagt gctacgagcg cttcagtcgc gagacagacg gggcgcagaa gcggcggatg 660
55 ctgcactgtg tgcagcgcgc gctgatccgc agccaggagc tgggcgacga gaagatccag 720
57 atcgtgagcc agatggtgga gctggtggag aaccgcacgc ggcaggtgga cagccacgtg 780
59 gagctgttcg aggcgcagca ggagctgggc gacacagcgg gcaacagcgg caaggctggc 840
61 gcggacaggc ccaaaggcga ggcggcagcg caggctgaca agcccaacag caagcgctca 900
63 cggcggcagc gcaacaacga gaaccgtgag aacgcgtcca gcaaccacga ccacgacgac 960
65 ggcgcctcgg gcacacccaa ggagaagaag gccaagacct ccaagaagaa gaagcgctcc 1020
67 aaggccaagg cggagcgaga ggcgtccgct gccgacctcc ccatcgaccc caacgaaccc 1080
69 acgtactgtc tgtgcaacca ggtctcctat ggggagatga tcggctgcga caacgacgag 1140
71 tgccccatcg agtggttcca cttctcgtgc gtggggctca atcataaacc caagggcaag 1200

RAW SEQUENCE LISTING DATE: 06/17/2002
PATENT APPLICATION: US/09/602,362C TIME: 13:47:25

Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF3\06172002\I602362C.raw

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     73 tggtactgtc ccaagtgccg gggggagaac gagaagacca tggacaaagc cctggagaaa 1260
     75 tccaaaaaag agagggctta caacaggtag tttgtggaca ggcgcctggt gtgaggagga 1320
     77 caaaataaac cgtgtattta ttacattgct gcctttgttg aggtgcaagg agtgtaaaat 1380
     79 gtatattttt aaagaatgtt agaaaaggaa ccattccttt catagggatg gcagtgattc 1440
     81 tgtttgcctt ttgttttcat tggtacacgt gtaacaagaa agtggtctgt ggatcagcat 1500
     83 tttagaaact acaaatatag gtttgattca aca                              1533
     86 <210> SEQ ID NO: 2
     87 <211> LENGTH: 1143
     88 <212> TYPE: DNA
     89 <213> ORGANISM: Homo sapiens
W--> 90 <400> SEQUENCE: 2
     91 gagtaacccg ataatatgcc gttgtccggc acggcgacga gaattcccag atatagcagt   60
     93 agcagtgatc ccgggcctgt ggctcggggc cggggctgca gttcggaccg cctcccgcga  120
     95 cccgcggggg ctcggagaca gtttcaggcc gcatctttgc tgacccgagg gtggggccgc  180
     97 gcgtggccgt ggaaacagat cctgaaggag ctagacgagt gctacgagcg cttcagtcgc  240
     99 gagacagacg gggcgcagaa gcggcggatg ctgcactgtg tgcagcgcgc gctgatccgc  300
    101 agccaggagc tgggcgacga gaagatccag atcgtgagcc agatggtgga gctggtggag  360
    103 aaccgcacgc ggcaggtgga cagccacgtg gagctgttcg aggcgcagca ggagctgggc  420
    105 gacacagtgg gcaacagcgg caaggttggc gcggacaggc ccaatggcga tgcggtagcg  480
    107 cagtctgaca agcccaacag caagcgctca cggcggcagc gcaacaacga gaaccgtgag  540
    109 aacgcgtcca gcaaccacga ccacgacgac ggcgcctcgg gcacacccaa ggagaagaag  600
    111 gccaagacct ccaagaagaa gaagcgctcc aaggccaagg cggagcgaga ggcgtcccct  660
    113 gccgacctcc ccatcgaccc caacgaaccc acgtactgtc tgtgcaacca ggtctcctat  720
    115 ggggagatga tcggctgcga caacgacgag tgccccatcg agtggttcca cttctcgtgc  780
    117 gtggggctca atcataaacc caagggcaag tggtactgtc ccaagtgccg gggggagaac  840
    119 gagaagacca tggacaaagc cctggagaaa tccaaaaaag agagggctta caacaggtag  900
    121 tttgtggaca ggcgcctggt gtgaggagga caaaataaac cgtgtattta ttacattgct  960
    123 gcctttgttg aggtgcaagg agtgtaaaat gtatattttt aaagaatgtt agaaaaggaa 1020
    125 ccattccttt catagggatg gcagtgattc tgtttgcctt ttgttttcat tggtacacgt 1080
    127 gtaacaagaa agtggtctgt ggatcagcat tttagaaact acaaatatag gtttgattca 1140
    129 aca                                                              1143
    132 <210> SEQ ID NO: 3
    133 <211> LENGTH: 742
    134 <212> TYPE: DNA
    135 <213> ORGANISM: Homo sapiens
    137 <400> SEQUENCE: 3
    138 cgccgtccac accccagcgg ccctgacgct gtcccctccg cgaccctcgc ctctggaaaa   60
    140 agtgacaggc aaggccacgc ccccgcgagg gccggcctcg agcccgcagc ccccagggcc  120
    142 tgggacgaga tcctgaagga gctagacgag tgctacgagc gcttcagtcg cgagacagac  180
    144 ggggcgcaga agcggcggat gctgcactgt gtgcagcgcg cgctgatccg cagccaggag  240
    146 ctgggcgacg agaagatcca gatcgtgagc cagatggtgg agctggtgga gaaccgcacg  300
    148 cggcaggtgg acagccacgt ggagctgttc gaggcgcagc aggagctggg cgacacagcg  360
    150 ggcaacagcg gcaaggctgg cgcggacagg cccaaaggcg aggcggcagc gcaggctgac  420
    152 aagcccaaca gcaagcgctc acggcggcag cgcaacaacg agaaccgtga gaacgcgtcc  480
    154 agcaaccacg accacgacga cggcgcctcg ggcacaccca aggagaagaa ggccaagacc  540
    156 tccaagaaga agaagcgctc caaggccaag gcggagcgag aggcgtcccc tgccgacctc  600
    158 cccatcgacc ccaacgaacc cacgtactgt ctgtgcaacc aggtctccta tggggagatg  660
    160 atcggctgcg acaacgacga gtgccccatc gagtggttcc acttctcgtg cgtggggctc  720
    162 aatcataaac ccaagggcaa gt                                         742
```

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/602,362C

DATE: 06/17/2002
TIME: 13:47:25

Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF3\06172002\I602362C.raw

165 <210> SEQ ID NO: 4
166 <211> LENGTH: 857
167 <212> TYPE: DNA
168 <213> ORGANISM: Homo sapiens
W--> 169 <400> SEQUENCE: 4
170 cctccgagaa cggtgtccat ggcacagggc gggaagagat aaggcctagg gaaggcgccc 60
172 ctcgggccta tccacctctt ctggggctcg gcactaggaa gcagcttccc tctcaggccc 120
174 ctttgtctcc aagccgttcc aaactgagta ccgggagacg acacaaaggg agggcggtga 180
176 cggatggcgc aggcgcggga gccgcctagg ctgctgggag tggtggtccg gccgcggaat 240
178 ggagatcctg aaggagctag acgagtgcta cgagcgcttc agtcgcgaga cagacggggc 300
180 gcagaagcgg cggatgctgc actgtgtgca gcgcgcgctg atccgcagcc aggagctggg 360
182 cgacgagaag atccagatcg tgagccagat ggtggagctg gtggagaacc gcacgcggca 420
184 ggtggacagc cacgtggagc tgttcgaggc gcagcaggag ctgggcgaca cagcgggcaa 480
186 cagcggcaag gctggcgcgg acaggcccaa aggcgaggcg gcagcgcagg ctgacaagcc 540
188 caacagcaag cgctcacggc ggcagcgcaa caacgagaac cgtgagaacg cgtccagcaa 600
190 ccacgaccac gacgacggcg cctcgggcac acccaaggag aagaaggcca agacctccaa 660
192 gaagaagaag cgctccaagg ccaaggcgga gcgagaggcg tcccctgccg acctccccat 720
194 cgaccccaac gaacccacgt actgtctgtg caaccaggtc tcctatgggg agatgatcgg 780
196 ctgcgacaac gacgagtgcc ccatcgagtg gttccacttc tcgtgcgtgg ggctcaatca 840
198 taaacccaag ggcaagt 857
201 <210> SEQ ID NO: 5
202 <211> LENGTH: 279
203 <212> TYPE: PRT
205 <213> ORGANISM: Homo sapiens
W--> 206 <400> SEQUENCE: 5
207 Met Leu Ser Pro Ala Asn Gly Glu Gln Leu His Leu Val Asn Tyr Val
208 1 5 10 15
210 Glu Asp Tyr Leu Asp Ser Ile Glu Ser Leu Pro Phe Asp Leu Gln Arg
211 20 25 30
213 Asn Val Ser Leu Met Arg Glu Ile Asp Ala Lys Tyr Gln Glu Ile Leu
214 35 40 45
216 Lys Glu Leu Asp Glu Cys Tyr Glu Arg Phe Ser Arg Glu Thr Asp Gly
217 50 55 60
219 Ala Gln Lys Arg Arg Met Leu His Cys Val Gln Arg Ala Leu Ile Arg
220 65 70 75 80
222 Ser Gln Glu Leu Gly Asp Glu Lys Ile Gln Ile Val Ser Gln Met Val
223 85 90 95
225 Glu Leu Val Glu Asn Arg Thr Arg Gln Val Asp Ser His Val Glu Leu
226 100 105 110
228 Phe Glu Ala Gln Gln Glu Leu Gly Asp Thr Val Gly Asn Ser Gly Lys
229 115 120 125
231 Val Gly Ala Asp Arg Pro Asn Gly Asp Ala Val Ala Gln Ser Asp Lys
232 130 135 140
234 Pro Asn Ser Lys Arg Ser Arg Arg Gln Arg Asn Asn Glu Asn Arg Glu
235 145 150 155 160
237 Asn Ala Ser Ser Asn His Asp His Asp Asp Gly Ala Ser Gly Thr Pro
238 165 170 175
240 Lys Glu Lys Lys Ala Lys Thr Ser Lys Lys Lys Lys Arg Ser Lys Ala
241 180 185 190

RAW SEQUENCE LISTING DATE: 06/17/2002
PATENT APPLICATION: US/09/602,362C TIME: 13:47:25

Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF3\06172002\I602362C.raw

```
243 Lys Ala Glu Arg Glu Ala Ser Pro Ala Asp Leu Pro Ile Asp Pro Asn
244         195                 200                 205
246 Glu Pro Thr Tyr Cys Leu Cys Asn Gln Val Ser Tyr Gly Glu Met Ile
247     210                 215                 220
249 Gly Cys Asp Asn Asp Glu Cys Pro Ile Glu Trp Phe His Phe Ser Cys
250 225                 230                 235                 240
252 Val Gly Leu Asn His Lys Pro Lys Gly Lys Trp Tyr Cys Pro Lys Cys
253                 245                 250                 255
255 Arg Gly Glu Asn Glu Lys Thr Met Asp Lys Ala Leu Glu Lys Ser Lys
256             260                 265                 270
258 Lys Glu Arg Ala Tyr Asn Arg
259         275
262 <210> SEQ ID NO: 6
263 <211> LENGTH: 210
264 <212> TYPE: PRT
265 <213> ORGANISM: Homo sapiens
W--> 266 <400> SEQUENCE: 6
267 Met Leu His Cys Val Gln Arg Ala Leu Ile Arg Ser Gln Glu Leu Gly
268 1               5                   10                  15
270 Asp Glu Lys Ile Gln Ile Val Ser Gln Met Val Glu Leu Val Glu Asn
271             20                  25                  30
273 Arg Thr Arg Gln Val Asp Ser His Val Glu Leu Phe Glu Ala Gln Gln
274         35                  40                  45
276 Glu Leu Gly Asp Thr Val Gly Asn Ser Gly Lys Val Gly Ala Asp Arg
277     50                  55                  60
279 Pro Asn Gly Asp Ala Val Ala Gln Ser Asp Lys Pro Asn Ser Lys Arg
280 65                  70                  75                  80
282 Ser Arg Arg Gln Arg Asn Asn Glu Asn Arg Glu Asn Ala Ser Ser Asn
283                 85                  90                  95
285 His Asp His Asp Asp Gly Ala Ser Gly Thr Pro Lys Glu Lys Lys Ala
286             100                 105                 110
288 Lys Thr Ser Lys Lys Lys Lys Arg Ser Lys Ala Lys Ala Glu Arg Glu
289         115                 120                 125
291 Ala Ser Pro Ala Asp Leu Pro Ile Asp Pro Asn Glu Pro Thr Tyr Cys
292     130                 135                 140
294 Leu Cys Asn Gln Val Ser Tyr Gly Glu Met Ile Gly Cys Asp Asn Asp
295 145                 150                 155                 160
297 Glu Cys Pro Ile Glu Trp Phe His Phe Ser Cys Val Gly Leu Asn His
298                 165                 170                 175
300 Lys Pro Lys Gly Lys Trp Tyr Cys Pro Lys Cys Arg Gly Glu Asn Glu
301             180                 185                 190
303 Lys Thr Met Asp Lys Ala Leu Glu Lys Ser Lys Lys Glu Arg Ala Tyr
304         195                 200                 205
306 Asn Arg
307     210
310 <210> SEQ ID NO: 7
311 <211> LENGTH: 235
312 <212> TYPE: PRT
313 <213> ORGANISM: Homo sapiens
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/602,362C

DATE: 06/17/2002
TIME: 13:47:25

Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF3\06172002\I602362C.raw

W--> 314 <400> SEQUENCE: 7
315 Met Glu Ile Leu Lys Glu Leu Asp Glu Cys Tyr Glu Arg Phe Ser Arg
316 1 5 10 15
318 Glu Thr Asp Gly Ala Gln Lys Arg Arg Met Leu His Cys Val Gln Arg
319 20 25 30
321 Ala Leu Ile Arg Ser Gln Glu Leu Gly Asp Glu Lys Ile Gln Ile Val
322 35 40 45
324 Ser Gln Met Val Glu Leu Val Glu Asn Arg Thr Arg Gln Val Asp Ser
325 50 55 60
327 His Val Glu Leu Phe Glu Ala Gln Gln Glu Leu Gly Asp Thr Val Gly
328 65 70 75 80
330 Asn Ser Gly Lys Val Gly Ala Asp Arg Pro Asn Gly Asp Ala Val Ala
331 85 90 95
333 Gln Ser Asp Lys Pro Asn Ser Lys Arg Ser Arg Arg Gln Arg Asn Asn
334 100 105 110
336 Glu Asn Arg Glu Asn Ala Ser Ser Asn His Asp His Asp Asp Gly Ala
337 115 120 125
339 Ser Gly Thr Pro Lys Glu Lys Lys Ala Lys Thr Ser Lys Lys Lys Lys
341 130 135 140
343 Arg Ser Lys Ala Lys Ala Glu Arg Glu Ala Ser Pro Ala Asp Leu Pro
344 145 150 155 160
346 Ile Asp Pro Asn Glu Pro Thr Tyr Cys Leu Cys Asn Gln Val Ser Tyr
347 165 170 175
349 Gly Glu Met Ile Gly Cys Asp Asn Asp Glu Cys Pro Ile Glu Trp Phe
350 180 185 190
352 His Phe Ser Cys Val Gly Leu Asn His Lys Pro Lys Gly Lys Trp Tyr
353 195 200 205
355 Cys Pro Lys Cys Arg Gly Glu Asn Glu Lys Thr Met Asp Lys Ala Leu
356 210 215 220
358 Glu Lys Ser Lys Lys Glu Arg Ala Tyr Asn Arg
359 225 230 235
362 <210> SEQ ID NO: 8
363 <211> LENGTH: 772
364 <212> TYPE: DNA
365 <213> ORGANISM: Homo sapiens
W--> 366 <220> FEATURE:
367 <221> NAME/KEY: CDS
368 <222> LOCATION: 695..714
W--> 369 <400> SEQUENCE: 8

n is found at 689, not 695 - must explain what residue n represents, See p. 6

370 aaagcgttct cggcggcagc gcaacaacta gaaccgtgag aacgcgtcca gcaaccgcga 60
372 cccacgacga cgtcacctcg ggcacgccca aggagaagaa agcccagacc tctaagaaga 120
374 agcagggctc catggccaag gcgtagcggc aggcgtcccc cgcagacctc cccatcgacc 180
376 ccagcgagcc ctcctactgg gagatgatcc gctgcgacaa cgaatgcccc atcgagtggt 240
378 tccgcttctc gtgtgtgagt ctcaaccata aaccaaagcg caagtggtac tgttccagat 300
380 gccggggaaa gaacgatggg caaagccctt gagaagtcca gaaaaaaaac agggcttata 360
382 acaggtagtt tggggacatg cgtctaatag tgaggagaac aaaataagcc agtgtgttga 420
384 ttacattgcc acctttgctg aggtgcagga agtgtaaaat gtatattttt aaagaatgtt 480
386 gttagaggcc gggcgcggtg gctcacgcct gtaatcccag cactttggga ggccgaggcg 540
388 gtcggatcac gaggtcagga gatcgagacc atcctggcta acacggtgaa accccgtctc 600

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 06/17/2002
PATENT APPLICATION: **US/09/602,362C** TIME: 13:47:26

Input Set : **A:\PTO.VSK.txt**
Output Set: **N:\CRF3\06172002\I602362C.raw**

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 235
Seq#:8; N Pos. 689,714
Seq#:15; N Pos. 1628,1752,1758,1769,1789,1873,1908,1915,1933,1970,1976,2022
Seq#:26; N Pos. 439,473,1789

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/602,362C

DATE: 06/17/2002
TIME: 13:47:26

Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF3\06172002\I602362C.raw

L:14 M:270 C: Current Application Number differs, Replaced Application Number
L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:29 M:283 W: Missing Blank Line separator, <220> field identifier
L:32 M:283 W: Missing Blank Line separator, <400> field identifier
L:39 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:1
L:39 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:180
L:90 M:283 W: Missing Blank Line separator, <400> field identifier
L:169 M:283 W: Missing Blank Line separator, <400> field identifier
L:206 M:283 W: Missing Blank Line separator, <400> field identifier
L:266 M:283 W: Missing Blank Line separator, <400> field identifier
L:314 M:283 W: Missing Blank Line separator, <400> field identifier
L:366 M:283 W: Missing Blank Line separator, <220> field identifier
L:369 M:283 W: Missing Blank Line separator, <400> field identifier
L:392 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:8
L:392 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:660
L:401 M:283 W: Missing Blank Line separator, <400> field identifier
L:419 M:283 W: Missing Blank Line separator, <400> field identifier
L:427 M:283 W: Missing Blank Line separator, <400> field identifier
L:435 M:283 W: Missing Blank Line separator, <400> field identifier
L:443 M:283 W: Missing Blank Line separator, <400> field identifier
L:451 M:283 W: Missing Blank Line separator, <220> field identifier
L:454 M:283 W: Missing Blank Line separator, <400> field identifier
L:509 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:15
L:509 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:1620
L:513 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:15
L:513 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:1740
L:517 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:15
L:517 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:1860
L:519 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:15
L:519 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:1920
L:521 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:15
L:521 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:1980
L:528 M:283 W: Missing Blank Line separator, <400> field identifier
L:633 M:283 W: Missing Blank Line separator, <400> field identifier
L:641 M:283 W: Missing Blank Line separator, <400> field identifier
L:649 M:283 W: Missing Blank Line separator, <400> field identifier
L:713 M:283 W: Missing Blank Line separator, <400> field identifier
L:720 M:283 W: Missing Blank Line separator, <400> field identifier
L:728 M:283 W: Missing Blank Line separator, <400> field identifier
L:872 M:283 W: Missing Blank Line separator, <400> field identifier
L:1132 M:283 W: Missing Blank Line separator, <400> field identifier
L:1139 M:283 W: Missing Blank Line separator, <400> field identifier
L:1147 M:283 W: Missing Blank Line separator, <220> field identifier
L:1150 M:283 W: Missing Blank Line separator, <400> field identifier
L:1165 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:26
L:1165 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:420
L:1209 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:26
L:1209 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:1740

VERIFICATION SUMMARY DATE: 06/17/2002
PATENT APPLICATION: **US/09/602,362C** TIME: 13:47:26

Input Set : **A:\PTO.VSK.txt**
Output Set: **N:\CRF3\06172002\I602362C.raw**

L:1280 M:283 W: Missing Blank Line separator, <400> field identifier
L:1479 M:283 W: Missing Blank Line separator, <400> field identifier
L:1487 M:283 W: Missing Blank Line separator, <400> field identifier